RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/627,685A_
Source: _1 Fw16_
Date Processed by STIC: _1/21/05_

ENTERED

RAW SEQUENCE LISTING　　　　　　　　DATE: 01/21/2005
PATENT APPLICATION:　**US/10/627,685A**　　　TIME: 16:03:09

Input Set : **A:\268.revised sequence.txt**
Output Set: **N:\CRF4\01212005\J627685A.raw**

```
 3 <110> APPLICANT: Cornell-Bell, Ann H.
 4         Pemberton, Karen E.
 5         Temple Jr., Davis L.
 6         Layer, Richard T.
 7         McCabe, R. Tyler
 8         Jones, Robert M.
 9         Cognetix, Inc.
11 <120> TITLE OF INVENTION: Uses of Kappa-Conotoxin PVIIA
13 <130> FILE REFERENCE: 2314-268
15 <140> CURRENT APPLICATION NUMBER: US 10/627,685A
16 <141> CURRENT FILING DATE: 2003-07-28
18 <150> PRIOR APPLICATION NUMBER: US 09/666,837
19 <151> PRIOR FILING DATE: 2000-09-21
21 <150> PRIOR APPLICATION NUMBER: US 60/219,438
22 <151> PRIOR FILING DATE: 2000-07-20
24 <150> PRIOR APPLICATION NUMBER: US 60/155,135
25 <151> PRIOR FILING DATE: 1999-09-22
27 <160> NUMBER OF SEQ ID NOS: 26
29 <170> SOFTWARE: PatentIn Ver. 2.0
31 <210> SEQ ID NO: 1
32 <211> LENGTH: 27
33 <212> TYPE: PRT
34 <213> ORGANISM: Conus purpurascens
36 <220> FEATURE:
37 <221> NAME/KEY: PEPTIDE
38 <222> LOCATION: (1)..(27)
39 <223> OTHER INFORMATION: Xaa at residue 2, 7, 18, 19, 22 and 25 may be Arg,
40         homoarginine, ornithine, Lys, N-methyl-Lys,
41         N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any
42         synthetic basic amino acid, His or halo-His; Xaa at
44 <220> FEATURE:
45 <221> NAME/KEY: PEPTIDE
46 <222> LOCATION: (1)..(27)
47 <223> OTHER INFORMATION: residue 4 may be Pro or Hyp; Xaa at residue 9 and
48         23 may be Phe,Tyr, meta-Tyr, ortho-Tyr, nor-Tyr,
49         mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
50         O-phospho-Tyr, nitro-Tyr, Trp (D or L), neo-Trp,
52 <220> FEATURE:
53 <221> NAME/KEY: PEPTIDE
54 <222> LOCATION: (1)..(27)
55 <223> OTHER INFORMATION: halo-Trp (D or L) or any synthetic aromatic amino
56         acid; Xaa at residue 11 is His or halo-His
58 <400> SEQUENCE: 1
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W--> 59 Cys Xaa Ile Xaa Asn Gln Xaa Cys Xaa Gln Xaa Leu Asp Asp Cys Cys
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     62 Ser Xaa Xaa Cys Asn Xaa Xaa Asn Xaa Cys Val
     63   20   25
     66 <210> SEQ ID NO: 2
     67 <211> LENGTH: 27
     68 <212> TYPE: PRT
     69 <213> ORGANISM: Conus purpurascens
     71 <220> FEATURE:
     72 <221> NAME/KEY: PEPTIDE
     73 <222> LOCATION: (1)..(27)
     74 <223> OTHER INFORMATION: Xaa is Hyp
     76 <400> SEQUENCE: 2
W--> 77 Cys Arg Ile Xaa Asn Gln Lys Cys Phe Gln His Leu Asp Asp Cys Cys
     78 1   5   10   15
     80 Ser Ala Lys Cys Asn Arg Phe Asn Lys Cys Val
     81   20   25
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     87 <213> ORGANISM: Conus purpurascens
     89 <220> FEATURE:
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     91 <222> LOCATION: (1)..(27)
     92 <223> OTHER INFORMATION: Xaa is Hyp
     94 <400> SEQUENCE: 3
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     96 1   5   10   15
     98 Ser Arg Lys Cys Asn Ala Phe Asn Lys Cys Val
     99   20   25
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    105 <213> ORGANISM: Conus purpurascens
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    108 <221> NAME/KEY: PEPTIDE
    109 <222> LOCATION: (1)..(27)
    110 <223> OTHER INFORMATION: Xaa is Hyp
    112 <400> SEQUENCE: 4
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    114 1   5   10   15
    116 Ser Arg Lys Cys Asn Arg Phe Asn Lys Cys Val
    117   20   25
    120 <210> SEQ ID NO: 5
    121 <211> LENGTH: 27
    122 <212> TYPE: PRT
    123 <213> ORGANISM: Conus purpurascens
    125 <220> FEATURE:
    126 <221> NAME/KEY: PEPTIDE

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    128 <223> OTHER INFORMATION: Xaa is Hyp
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    135             20                  25
    138 <210> SEQ ID NO: 6
    139 <211> LENGTH: 27
    140 <212> TYPE: PRT
    141 <213> ORGANISM: Conus purpurascens
    143 <220> FEATURE:
    144 <221> NAME/KEY: PEPTIDE
    145 <222> LOCATION: (1)..(27)
    146 <223> OTHER INFORMATION: Xaa is Hyp
    148 <400> SEQUENCE: 6
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    150   1               5                  10                  15
    152 Ser Arg Lys Cys Asn Arg Phe Asn Lys Cys Val
    153             20                  25
    156 <210> SEQ ID NO: 7
    157 <211> LENGTH: 27
    158 <212> TYPE: PRT
    159 <213> ORGANISM: Conus purpurascens
    161 <220> FEATURE:
    162 <221> NAME/KEY: PEPTIDE
    163 <222> LOCATION: (1)..(27)
    164 <223> OTHER INFORMATION: Xaa is Hyp
    166 <400> SEQUENCE: 7
W-->167 Cys Arg Ile Xaa Asn Gln Lys Cys Ala Gln His Leu Asp Asp Cys Cys
    168   1               5                  10                  15
    170 Ser Arg Lys Cys Asn Arg Phe Asn Lys Cys Val
    171             20                  25
    174 <210> SEQ ID NO: 8
    175 <211> LENGTH: 27
    176 <212> TYPE: PRT
    177 <213> ORGANISM: Conus purpurascens
    179 <220> FEATURE:
    180 <221> NAME/KEY: PEPTIDE
    181 <222> LOCATION: (1)..(27)
    182 <223> OTHER INFORMATION: Xaa is Hyp
    184 <400> SEQUENCE: 8
W-->185 Cys Arg Ile Xaa Asn Gln Lys Cys Phe Gln His Leu Asp Asp Cys Cys
    186   1               5                  10                  15
    188 Ser Arg Lys Cys Asn Arg Phe Asn Ala Cys Val
    189             20                  25
    192 <210> SEQ ID NO: 9
    193 <211> LENGTH: 27
    194 <212> TYPE: PRT
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PATENT APPLICATION: US/10/627,685A   TIME: 16:03:09

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    197 <220> FEATURE:
    198 <221> NAME/KEY: PEPTIDE
    199 <222> LOCATION: (1)..(27)
    200 <223> OTHER INFORMATION: Xaa is Hyp
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    204   1               5                   10                  15
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    207               20                  25
    210 <210> SEQ ID NO: 10
    211 <211> LENGTH: 27
    212 <212> TYPE: PRT
    213 <213> ORGANISM: Conus purpurascens
    215 <220> FEATURE:
    216 <221> NAME/KEY: PEPTIDE
    217 <222> LOCATION: (1)..(27)
    218 <223> OTHER INFORMATION: Xaa is Hyp
    220 <400> SEQUENCE: 10
W--> 221 Cys Arg Ile Xaa Asn Gln Ala Cys Phe Gln His Leu Asp Asp Cys Cys
    222   1               5                   10                  15
    224 Ser Arg Lys Cys Asn Arg Phe Asn Lys Cys Val
    225               20                  25
    228 <210> SEQ ID NO: 11
    229 <211> LENGTH: 27
    230 <212> TYPE: PRT
    231 <213> ORGANISM: Conus purpurascens
    233 <220> FEATURE:
    234 <221> NAME/KEY: PEPTIDE
    235 <222> LOCATION: (1)..(27)
    236 <223> OTHER INFORMATION: Xaa is Hyp
    238 <400> SEQUENCE: 11
W--> 239 Cys Arg Ile Xaa Asn Gln Lys Cys Met Gln His Leu Asp Asp Cys Cys
    240   1               5                   10                  15
    242 Ser Arg Lys Cys Asn Arg Phe Asn Lys Cys Val
    243               20                  25
    245 <210> SEQ ID NO: 12
    246 <211> LENGTH: 27
    247 <212> TYPE: PRT
    248 <213> ORGANISM: Conus purpurascens
    250 <220> FEATURE:
    251 <221> NAME/KEY: PEPTIDE
    252 <222> LOCATION: (1)..(27)
    253 <223> OTHER INFORMATION: Xaa is Hyp
    255 <400> SEQUENCE: 12
W--> 256 Cys Arg Ile Xaa Asn Gln Lys Cys Tyr Gln His Leu Asp Asp Cys Cys
    257   1               5                   10                  15
    259 Ser Arg Lys Cys Asn Arg Phe Asn Lys Cys Val
    260               20                  25
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263 <211> LENGTH: 27
264 <212> TYPE: PRT
265 <213> ORGANISM: Conus purpurascens
267 <220> FEATURE:
268 <221> NAME/KEY: PEPTIDE
269 <222> LOCATION: (1)..(27)
270 <223> OTHER INFORMATION: Xaa is Hyp
272 <400> SEQUENCE: 13
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276 Ser Arg Lys Cys Asn Arg Phe Asn Lys Cys Val
277              20                  25
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282 <212> TYPE: PRT
283 <213> ORGANISM: Conus purpurascens
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287 <222> LOCATION: (1)..(27)
288 <223> OTHER INFORMATION: Xaa is Hyp
290 <400> SEQUENCE: 14
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294 Ser Arg Lys Cys Asn Arg Phe Asn Lys Cys Val
295              20                  25
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299 <211> LENGTH: 27
300 <212> TYPE: PRT
301 <213> ORGANISM: Conus purpurascens
303 <220> FEATURE:
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305 <222> LOCATION: (1)..(27)
306 <223> OTHER INFORMATION: Xaa is Hyp
308 <400> SEQUENCE: 15
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313              20                  25
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317 <211> LENGTH: 27
318 <212> TYPE: PRT
319 <213> ORGANISM: Conus purpurascens
321 <220> FEATURE:
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323 <222> LOCATION: (1)..(27)
324 <223> OTHER INFORMATION: Xaa is Hyp
326 <400> SEQUENCE: 16
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 2,4,7,9,11,16,19,22,23,25
Seq#:2; Xaa Pos. 4
Seq#:3; Xaa Pos. 4
Seq#:4; Xaa Pos. 4
Seq#:5; Xaa Pos. 4
Seq#:6; Xaa Pos. 4
Seq#:7; Xaa Pos. 4
Seq#:8; Xaa Pos. 4
Seq#:9; Xaa Pos. 4
Seq#:10; Xaa Pos. 4
Seq#:11; Xaa Pos. 4
Seq#:12; Xaa Pos. 4
Seq#:13; Xaa Pos. 4
Seq#:14; Xaa Pos. 4
Seq#:15; Xaa Pos. 4
Seq#:16; Xaa Pos. 4
Seq#:17; Xaa Pos. 4
Seq#:18; Xaa Pos. 4
Seq#:19; Xaa Pos. 4
Seq#:20; Xaa Pos. 4
Seq#:21; Xaa Pos. 4
Seq#:22; Xaa Pos. 4
Seq#:23; Xaa Pos. 4
Seq#:25; Xaa Pos. 4
Seq#:26; Xaa Pos. 4

VERIFICATION SUMMARY   DATE: 01/21/2005
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Input Set : **A:\268.revised sequence.txt**
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```
L:59 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
M:341 Repeated in SeqNo=1
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:95 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:167 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:291 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:327 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:345 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:363 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:381 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:417 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:483 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:501 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
```